

```

<!--StartFragment-->RESULT 1
US-09-828-217-1
; Sequence 1, Application US/09828217
; Patent No. US20010051710A1
;   GENERAL INFORMATION:
;       APPLICANT: HIRANO, TOSHIO
;               KAISHO, TSUNEFASU
;       TITLE OF INVENTION: MEMBRANE PROTEIN POLYPEPTIDE HAVING
;               PRE-B CELL GROWTH-SUPPORTING ABILITY AND A GENE THEREOF
;       NUMBER OF SEQUENCES: 2
;       CORRESPONDENCE ADDRESS:
;               ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
;               P.C.
;               STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
;               CITY: ARLINGTON
;               STATE: VA
;               COUNTRY: USA
;               ZIP: 22202
;       COMPUTER READABLE FORM:
;               MEDIUM TYPE: Floppy disk
;               COMPUTER: IBM PC compatible
;               OPERATING SYSTEM: PC-DOS/MS-DOS
;               SOFTWARE: PatentIn Release #1.0, Version #1.30
;       CURRENT APPLICATION DATA:
;               APPLICATION NUMBER: US/09/828,217
;               FILING DATE: 09-Apr-2001
;               CLASSIFICATION: <Unknown>
;       PRIOR APPLICATION DATA:
;               APPLICATION NUMBER: 09/182,563
;               FILING DATE: <Unknown>
;               APPLICATION NUMBER: PCT/JP94/01732
;               FILING DATE: 14-OCT-1994
;               APPLICATION NUMBER: JP 5-281622
;               FILING DATE: 15-OCT-1993
;       ATTORNEY/AGENT INFORMATION:
;               NAME: OBLON, NORMAN F.
;               REGISTRATION NUMBER: 24,618
;               REFERENCE/DOCKET NUMBER: 7625-001-0 PCT
;       TELECOMMUNICATION INFORMATION:
;               TELEPHONE: 703-413-3000
;               TELEFAX: 703-413-2220
;       INFORMATION FOR SEQ ID NO: 1:
;       SEQUENCE CHARACTERISTICS:
;               LENGTH: 180 amino acids
;               TYPE: amino acid
;               STRANDEDNESS: single
;               TOPOLOGY: linear
;       MOLECULE TYPE: peptide
;       SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-828-217-1

```

```

Query Match          100.0%;  Score 889;  DB 3;  Length 180;
Best Local Similarity 100.0%;  Pred. No. 2.9e-77;
Matches 180;  Conservative 0;  Mismatches 0;  Indels 0;  Gaps 0;

```

```

Qy      1 MASTSYDYCRVPMEDGDKRCKLLLGIGILVLLIIVILGVPLIIFTIKANSEACRDGLRAV 60
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1 MASTSYDYCRVPMEDGDKRCKLLLGIGILVLLIIVILGVPLIIFTIKANSEACRDGLRAV 60

Qy      61 MECRNVTLLQQLTEAQKGFQDVEAQAATCNHTVMALMASLDAEKAQGQKKVEELEGEI 120
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||

```

Db . 61 MECRNVTHLLQQELTEAQKGFQDVEAQAATCNHTVMALMASLDAEKAQGQKKVEELEGEI 120

Qy 121 TTLNHKLQDASAEVERLRRENQVLSVRIADKKYYPSSQDSSSAAAPQLLIVLLGLSALLQ 180

Db 121 TTLNHKLQDASAEVERLRRENQVLSVRIADKKYYPSSQDSSSAAAPQLLIVLLGLSALLQ 180

<!--EndFragment-->